

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: June 24, 2002, 20:48:36 ; Search time 16.32 Seconds
(without alignments) 382.709 Million cell updates/sec

Title: US-09-664-326-23

Perfect score: 368
Sequence: 1 LHYTCTESGQNLCLCEGSN.....PKQSHNDGFEEIPEEYIQ 65

Scoring table: BLOSUM62
Gapop 10.0 , Gapect 0.5

Searched: 283138 seqs, 96089334 residues

Total number of hits satisfying chosen parameters: 283138

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Database :

1: PIR_71:*
2: PIR1:*
3: PIR2:*
4: PIR3:*
5: PIR4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	360	97.8	65	1 HULXH	thrombin inhibitor
2	353	95.9	65	2 S05674	hirudin IIB - medi
3	352	95.7	65	3 S78521	hirudin IIB - me
4	350	95.1	65	2 S05673	hirudin IIA - medi
5	348	94.6	65	2 S05676	hirudin IIR - medi
6	348	94.6	65	2 S05678	hirudin IIR - medi
7	346	94.0	65	2 S78520	hirudin IIR - me
8	345	93.8	65	2 S05677	hirudin IIR - me
9	345	93.8	65	2 S05675	hirudin IIR - me
10	343	93.2	65	2 S05679	hirudin IIR - medi
11	333	90.5	72	2 A37417	thrombin inhibitor
12	315	85.6	66	2 A24350	thrombin inhibitor
13	288	78.3	55	2 S05672	hirudin I - medi
14	249	67.7	63	2 A53883	hirudin I - medi
15	247	67.1	84	2 S33329	hirudin HM2 - leec
16	230	62.5	84	2 S33328	hirudin HM1 - leec
17	215	58.4	63	1 A42207	hirudin P6 - leech
18	173	47.0	62	1 HULXLM	hirudin P18 - lee
19	90	24.5	17	2 S05671	hirudin Ia - medic
20	74	20.1	23	2 T30201	Notch homolog prot
21	69	18.8	17	2 A42125	trophozoite cystei
22	68.5	18.6	1071	1 PEXBYVA	H+-transporting AT
23	68.5	18.6	2233	2 T28659	surface protein 51
24	66	17.9	761	2 T09052	hypothetical prote
25	66	17.9	2150	2 T32497	hypothetical prote
26	66	17.9	5376	2 T42215	zonadhesin - mouse
27	65	17.7	558	2 T15448	hypothetical prote
28	65	17.7	1043	2 T19734	hypothetical prote
29	65	17.7	1661	2 T31330	head-activator bin

30	64	17.4	317	2	I46916	insulin-like growt
31	64	17.4	1743	2	T26859	hypothetical prote
32	63.5	17.3	318	2	A82319	glutathione synthe
33	63.5	17.3	474	2	S18452	variant surface gl
34	63.5	17.3	755	2	A44315	collagen oligomer
35	63.5	17.3	1722	2	E89753	protein F11C7.4 [1
36	63.5	17.3	2321	2	S78549	notch protein - h
37	63.5	17.3	2703	1	A24420	notch protein - fr
38	63	17.1	80	2	T10183	antifungal protein
39	63	17.1	778	2	T05341	S-receptor kinase
40	62.5	17.0	513	1	Q01486	activin receptor I
41	62.5	17.0	1820	2	A55494	latent transformin
42	62	16.8	79	2	T10243	antifungal protein
43	62	16.8	79	2	T07917	antifungal protein
44	62	16.8	80	2	T02621	probable antifunga
45	62	16.8	80	2	T02622	probable antifunga

ALIGNMENTS

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RESULT 1
HULXH
thrombin inhibitor (hirudin) - medicinal leech
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 30-Nov-1980 #sequence:revision 03-Aug-1984 #text_change 07-May-1999
C:Accession: A91318; A94429; A60811; A01289
R:Dot, J., Muller, H.P.; Seemuller, U.; Chang, J.Y.
FEBS Lett. 165, 180-183, 1984
A:Title: The complete amino acid sequence of hirudin, a thrombin specific inhibitor.
A:Reference number: A91318
A:Accession: A91318
A:Molecule type: protein
A:Residues: 1-65 <DOD>
R:Peterson, T.E.; Roberts, H.R.; Sottrup-Jensen, L.; Magnusson, S.; Bagdy, D.
Anal. Biochem. 161, 514-518, 1987
A:Title: Rapid purification and revised amino-terminal sequence of hirudin: a specific
A:Reference number: A60811; MUID:87211066
A:Accession: A60811
A:Molecule type: protein
A:Residues: 1-65 <DOD>
R:Mao, S.J.T.; Yates, M.T.; Blankenship, D.T.; Cardin, A.D.; Krstenansky, J.L.; Loven
Anal. Biochem. 161, 514-518, 1987
A:Title: Rapid purification and revised amino-terminal sequence of hirudin: a specific
A:Reference number: A60811; MUID:87211066
A:Accession: A60811
A:Molecule type: protein
A:Residues: 1-32, 'N', 34-43 <MAO>
A>Note: the authors suggest that their identification of 33-Asn is correct and that 3
present a natural variant of hirudin
C:Comment: Hirudin is a potent thrombin-specific protease inhibitor.
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; sulfitoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status experimental
F:63/Binding site: sulfate (Tyr) (covalent) #status experimental

Query Match 97.8%; Score 360; DB 1; Length 65;
Best local similarity 96.9%; Pred. No. 7.8e-30;
Matches 63; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 LHYTCTESGQNLCLCEGSNNVCGGKNCITLGSDEKNCVYEGTPKQSHNDGFEEIP 60
DB 1 VYITCTESGQNLCLCEGSNNVCGGKNCITLGSDEKNCVYEGTPKQSHNDGFEEIP 60
QY 61 EYLIQ 65
DB 61 EYLIQ 65

RESULT 2
S05674
hirudin IIB - medicinal leech
N:Alternate names: thrombin inhibitor

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C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 17-Jul-1998
C:Accession: S05674
R:Scharf, M.; Engels, J.; Triplier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945
A:Accession: S05674
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.9%; Score 353; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 4e-29;
Matches 61; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

QY 1 LVTYDCTESGQNLCLCEGSNVCQGQNKCTILGSDGKNCVGTGCTPKPQSHNDGDFEETP 60
:|||||
Db 1 ITVTDCTESGQNLCLCEGSNVCQGNKCTILGNGEENOCVTGCTPKPQSHNDGDFEETP 60

QY 61 EEYLIQ 65
|||||
Db 61 EEYLIQ 65

RESULT 3
S78521
hirudin IIIB' - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
C:Accession: S78521
R:Scharf, M.; Engels, J.; Triplier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945
A:Accession: S78521
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.7%; Score 352; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 5.1e-29;
Matches 61; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 LVTYDCTESGQNLCLCEGSNVCQGQNKCTILGSDGKNCVGTGCTPKPQSHNDGDFEETP 60
:|||||
Db 1 VVTYDCTESGQNLCLCEGSNVCQGNKCTILGNGEENOCVTGCTPKPQSHNDGDFEETP 60

QY 61 EEYLIQ 65
|||||
Db 61 EEYLIQ 65

RESULT 4
S05673
hirudin IIA - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05673
R:Scharf, M.; Engels, J.; Triplier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945

A:Accession: S05673
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:47/inhibitory site: Lys (thrombin) #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 95.1%; Score 350; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 8.1e-29;
Matches 61; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTYDCTESGQNLCLCEGSNVCQGQNKCTILGSDGKNCVGTGCTPKPQSHNDGDFEETP 60
:|||||
Db 1 ITVTDCTESGQNLCLCEGSNVCQGNKCTILGNGEENOCVTGCTPKPQSHNDGDFEETP 60

QY 61 EEYLIQ 65
|||||
Db 61 EEYLIQ 65

RESULT 5
S05676
hirudin III - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05676
R:Scharf, M.; Engels, J.; Triplier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945
A:Accession: S05676
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted
F:63/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 94.6%; Score 348; DB 2; Length 65;
Best Local Similarity 93.8%; Pred. No. 1.3e-28;
Matches 61; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTYDCTESGQNLCLCEGSNVCQGQNKCTILGSDGKNCVGTGCTPKPQSHNDGDFEETP 60
:|||||
Db 1 VVTYDCTESGQNLCLCEGSNVCQGNKCTILGNGEENOCVTGCTPKPQSHNDGDFEETP 60

QY 61 EEYLIQ 65
|||||
Db 61 EEYLIQ 65

RESULT 6
S05678
hirudin II - medicinal leech
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 17-Jul-1998
C:Accession: S05678
R:Scharf, M.; Engels, J.; Triplier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.
A:Reference number: S05671; MUID:90005945
A:Accession: S05678
A:Molecule type: protein
A:Residues: 1-65 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F:6-14,16-28,22-39/Disulfide bonds: #status predicted

RESULT 11
A37417
thrombin inhibitor (hirudin) type II precursor - medicinal leech (fragment)
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 28-Apr-1993 #sequence_revision 28-Apr-1993 #text_change 16-Jul-1999
C:Accession: A37417
R:Harvey, R.P.; Degryse, E.; Stefan, L.; Schamber, F.; Cazenave, J.P.; Courtney, M.; To
Proc. Natl. Acad. Sci. U.S.A. 83, 1084-1088, 1986
A:Title: Cloning and expression of a cDNA coding for the anticoagulant hirudin from the
A:Reference number: A37417; MUID:86149219
A:Accession: A37417
A:Status: preliminary; not compared with conceptual translation
A:Molecule type: mRNA
A:Residues: 1-72 <HAR>
A:Cross-references: GB:M12693; NID:g159224; PIDN:AAA29195.1; PID:g159225
C:Superfamily: thrombin inhibitor

Query Match 90.5%; Score 333; DB 2; Length 72;
Best Local Similarity 87.7%; Pred. No. 4.7e-27;
Matches 57; Conservative 6; Mismatches 2; Indels 0; Gaps 0;

QY 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 60
:|||||
DB 8 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 67
QY 61 EEYIQ 65
:|||||
DB 68 EEYIQ 72

RESULT 12
A24350
thrombin inhibitor (hirudin PA) - medicinal leech
N:Alternate names: hirudin (plasminogen activator-type)
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 18-Jun-1993
C:Accession: A24350
R:Dodd, J.; Machleidt, W.; Seemüller, U.; Maschler, R.; Fritzt, H.
Biol. Chem. Hoppe-Seyler 367, 803-811, 1986
A:Title: Isolation and characterization of hirudin isolinhibitors and sequence analysis
A:Reference number: A24350; MUID:87026247
A:Accession: A24350
A:Molecule type: protein
A:Residues: 1-66 <DOD>
C:Superfamily: thrombin inhibitor

Query Match 85.6%; Score 315; DB 2; Length 66;
Best Local Similarity 87.1%; Pred. No. 2.9e-25;
Matches 54; Conservative 4; Mismatches 4; Indels 0; Gaps 0;

QY 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 60
:|||||
DB 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 60
QY 61 EE 62
:|:
DB 61 ED 62

RESULT 13
S05672
hirudin I - medicinal leech (fragments)
N:Alternate names: thrombin inhibitor
C:Species: Hirudo medicinalis (medicinal leech)
C:Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 17-Jul-1998
C:Accession: S05672
R:Scharf, M.; Engels, J.; Tripier, D.
FEBS Lett. 255, 105-110, 1989
A:Title: Primary structures of new 'iso-hirudins'.

A:Reference number: S05671; MUID:90005945
A:Accession: S05672
A:Molecule type: Protein
A:Residues: 1-55 <SCH>
C:Superfamily: thrombin inhibitor
C:Keywords: anticoagulant; serine proteinase inhibitor; sulfoprotein
F/6-14/Disulfide bonds: #status predicted
F/53/Binding site: sulfate (Tyr) (covalent) #status predicted

Query Match 78.3%; Score 288; DB 2; Length 55;
Best Local Similarity 81.5%; Pred. No. 1.3e-22;
Matches 53; Conservative 1; Mismatches 1; Indels 10; Gaps 1;

QY 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 60
:|||||
DB 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 50
QY 61 EEYIQ 65
:|||||
DB 51 EEYIQ 55

RESULT 14
A53883
hirudin HVI homolog bufrudin - leech (Hirudinaria manillensis)
C:Species: Hirudinaria manillensis
C:Date: 27-Sep-1994 #sequence_revision 18-Nov-1994 #text_change 09-Mar-1996
C:Accession: A53883
R:Electricwala, A.; Hartwell, R.; Scawen, M.D.; Atkinson, T.
J. Protein Chem. 12, 365-370, 1993
A:Title: The complete amino acid sequence of a hirudin variant from the leech Hirudin
A:Reference number: A53883; MUID:94000343
A:Accession: A53883
A:Status: preliminary
A:Molecule type: protein
A:Residues: 1-63 <ELE>
A:Experimental source: head portions
A:Note: sequence extracted from NCBI backbone (NCBI:139162)
C:Superfamily: thrombin inhibitor

Query Match 67.7%; Score 249; DB 2; Length 63;
Best Local Similarity 67.7%; Pred. No. 1.3e-18;
Matches 44; Conservative 8; Mismatches 11; Indels 2; Gaps 1;

QY 1 LVTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 60
:|||||
DB 1 VSTDTESGQNLCEGSSNVCQGKNCILGSDGKNCVTEGTPKPSHNDGDFEETP 58
QY 61 EEYIQ 65
:|:
DB 59 DEXIK 63

RESULT 15
S33329
hirudin HM2 - leech (Hirudinaria manillensis)
C:Species: Hirudinaria manillensis
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Jul-1999
C:Accession: S33329
R:Scacchi, E.; Nitti, G.; Valassina, B.; Orsini, G.; Visco, C.; Ferreira, M.; Sawyer,
Eur. J. Biochem. 214, 295-304, 1993
A:Title: Novel hirudin variants from the leech Hirudinaria manillensis. Amino acid se
A:Reference number: S33328; MUID:93285156
A:Accession: S33329
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-84 <SCA>
A:Cross-references: EMBL:X72786; NID:g312674; PIDN:CAA51293.1; PID:g312675
C:Genetics:
A:Introns: 21/1; 37/3; 61/1
C:Superfamily: thrombin inhibitor

